

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/591, 725
Source: IFWP
Date Processed by STIC: 09/13/2006

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 09/13/2006

PATENT APPLICATION: US/10/591,725

TIME: 11:08:53

Input Set : A:\021080002U4.txt

Output Set: N:\CRF4\09132006\J591725.raw

```

4 <110> APPLICANT: Leonard, Joan
6 <120> TITLE OF INVENTION: CHICKEN ANEMIA VIRUS VACCINE FROM CELL
7 LINE
9 <130> FILE REFERENCE: 02108.0002U4
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/591,725
12 <141> CURRENT FILING DATE: 2006-09-05
14 <150> PRIOR APPLICATION NUMBER: PCT/US05/07148
15 <151> PRIOR FILING DATE: 2005-03-07
17 <150> PRIOR APPLICATION NUMBER: 60/550,573
18 <151> PRIOR FILING DATE: 2004-03-05
20 <160> NUMBER OF SEQ ID NOS: 2
22 <170> SOFTWARE: FastSEQ for Windows Version 4.0
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 2294
26 <212> TYPE: DNA
27 <213> ORGANISM: Chicken anemia virus
29 <400> SEQUENCE: 1
30 gcatccgag tggttactat tccatcacca ttctagcctg tacacagaaa gtcaagatgg 60
31 acgaatcgct cgacttcgct cgcgattcgt cgaaggcggg gggccggagg ccccccgggtg 120
32 gccccctcc aacgagtggg gcatgtacag ggggggtacgt catcagtaca ggggggtacg 180
33 tcacaaaaag gcggtcccggt acaggggggt acgtcacgag tacagggggg tacgtcacag 240
34 ccaatcagaa gctgccacgt tgcgaaagtg acgtttcgaa aatgggcggc gcaagcctct 300
35 ctatatattg agcgcacata ccggtcggca gtaggtatac gcaaggcggg ccgggtggat 360
36 gcacgggaac cccggaacac cggccgctgg gggcagtgaa tcggcgctta gccgagaggg 420
37 gcagcctggg gccagcggag ccgcgcaggg gcaagtaatt tcaaatgaac gctctccaag 480
38 aagatactcc acccggacca tcaacggtgt tcaggccacc aacaagttca cggccgttgg 540
39 aaaccctca ctgcagagag atccggattg gtatcgctgg aattacaatc actctatcgc 600
40 tgtgtggctg cgcgaaatgct cgcgctccca cgctaagatc tgcaactgag gacaattcag 660
41 aaagcactgg tttcaagaat gtgccggact tgaggaccga tcaaccgaag cctccctcga 720
42 agaagcgatc ctgcgacccc tccgagtaca gggtaagcga gctaaaagaa agcttgatta 780
43 ccactactcc cagccgaccc cgaaccgcaa gaaggtgtat aagactgtaa gatggcaaga 840
44 cgagctcgca gaccgagagg ccgattttac gccttcagaa gaggacgggtg gcaccacctc 900
45 aagcgacttc gacgaagata taaatttcga catcggagga gacagcggta tcgtagacga 960
46 gcttttagga aggcctttca caacccccgc cccggtacgt atagtgtgag gctgccgaac 1020
47 cccaatcta ccatgactat ccgcttccaa ggagtcattt ttctcacaga aggactcatt 1080
48 ctgcctaaaa acagcacagc ggggggctat gcagaccaca tgtacggggc gagagtcgcc 1140
49 aagatctctg tgaacctgaa agagttcctg ctagecgtcaa tgaacctgac atacgtgagc 1200
50 aaaatcggag gccccatcgc cgggtgagttg attgcggacg ggtctaaatc acaagccgcg 1260
51 gagaactggc ctaattgctg gctgccgcta gataataacg tgccctccgc gacaccatcg 1320
52 gcatggtgga gatgggcctt aatgatgatg cagcccacgg actcttgccg gttctttaat 1380
53 caccctaagc agatgacctt gcaagacatg ggtcgcattg ttgggggctg gcacctatct 1440
54 cgacacattg aaaccgcctt tcagctcctt gccactaaga atgagggatc cttcagcccc 1500
55 gtggcgagtc ttctctccca gggagagtag ctcacgcgtc gggacgatgt taaatacagc 1560

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56 agcgatcacc agaaccggtg gcgaaaaggc gaacaaccga tgacggggggg tattgcttat      1620
57 gcgaccggga aaatgagacc cgacgagcaa cagtaccctg ctatgcccc agaccccccg      1680
58 atcatcacca gtactacagc gcaaggcacg caagtccgct gcatgaatag cacgcaagct      1740
59 tgggtggtcat gggacacata tatgagcttt gcaacactca cagcactcgg tgcacaatgg      1800
60 tcttttcctc cagggcaacg ttcagtttct agacggtcct tcaaccacca caaggcgaga      1860
61 ggagccgggg accccaaagg ccagagatgg cacaccctgg tgccgctcgg cacggagacc      1920
62 atcaccgaca gctacatggg agcaccgcga tcagagctgg acacaaattt ctttacgctt      1980
63 tacgtagcgc aaggtacaaa taagtcgcag cagtacaagt tcggcacagc tacatacgcg      2040
64 ctaaaggagc cggtaatgaa gacgatttca tgggcagtgg tacgcgtcca gtcggtctgg      2100
65 caactgggta acaggcagag accataccca tgggacgtca actgggcca cagcaccatg      2160
66 tactggggcg ggcagccctg aaaagggggg gggctaaagc ccccccttga accccccctt      2220
67 ggggggggatt cccccccaga ccccccttt aaatagcact caataaacgc agcaattggc      2280
68 tttatcgcac aatc                                     2294
70 <210> SEQ ID NO: 2
71 <211> LENGTH: 786
72 <212> TYPE: PRT
73 <213> ORGANISM: Chicken anemia virus
75 <400> SEQUENCE: 2
76 Met Ala Arg Arg Ala Arg Arg Pro Arg Gly Arg Phe Tyr Ala Phe Arg
77 1 5 10 15
78 Arg Gly Arg Trp His His Leu Lys Arg Leu Arg Arg Arg Tyr Lys Phe
79 20 25 30
80 Arg His Arg Arg Arg Gln Arg Tyr Arg Arg Arg Ala Phe Arg Lys Ala
81 35 40 45
82 Phe His Asn Pro Arg Pro Gly Thr Tyr Ser Val Arg Leu Pro Asn Pro
83 50 55 60
84 Gln Ser Thr Met Thr Ile Arg Phe Gln Gly Val Ile Phe Leu Thr Glu
85 65 70 75 80
86 Gly Leu Ile Leu Pro Lys Asn Ser Thr Ala Gly Gly Tyr Ala Asp His
87 85 90 95
88 Met Tyr Gly Ala Arg Val Ala Lys Ile Ser Val Asn Leu Lys Glu Phe
89 100 105 110
90 Leu Leu Ala Ser Met Asn Leu Thr Tyr Val Ser Lys Ile Gly Gly Pro
91 115 120 125
92 Ile Ala Gly Glu Leu Ile Ala Asp Gly Ser Lys Ser Gln Ala Ala Glu
93 130 135 140
94 Asn Trp Pro Asn Cys Trp Leu Pro Leu Asp Asn Asn Val Pro Ser Ala
95 145 150 155 160
96 Thr Pro Ser Ala Trp Trp Arg Trp Ala Leu Met Met Met Gln Pro Thr
97 165 170 175
98 Asp Ser Cys Arg Phe Phe Asn His Pro Lys Gln Met Thr Leu Gln Asp
99 180 185 190
100 Met Gly Arg Met Phe Gly Gly Trp His Leu Phe Arg His Ile Glu Thr
101 195 200 205
102 Arg Phe Gln Leu Leu Ala Thr Lys Asn Glu Gly Ser Phe Ser Pro Val
103 210 215 220
104 Ala Ser Leu Leu Ser Gln Gly Glu Tyr Leu Thr Arg Arg Asp Asp Val
105 225 230 235 240
106 Lys Tyr Ser Ser Asp His Gln Asn Arg Trp Arg Lys Gly Glu Gln Pro

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107				245				250				255				
108	Met	Thr	Gly	Gly	Ile	Ala	Tyr	Ala	Thr	Gly	Lys	Met	Arg	Pro	Asp	Glu
109				260				265						270		
110	Gln	Gln	Tyr	Pro	Ala	Met	Pro	Pro	Asp	Pro	Pro	Ile	Ile	Thr	Ser	Thr
111			275					280					285			
112	Thr	Ala	Gln	Gly	Thr	Gln	Val	Arg	Cys	Met	Asn	Ser	Thr	Gln	Ala	Trp
113		290					295					300				
114	Trp	Ser	Trp	Asp	Thr	Tyr	Met	Ser	Phe	Ala	Thr	Leu	Thr	Ala	Leu	Gly
115	305					310					315				320	
116	Ala	Gln	Trp	Ser	Phe	Pro	Pro	Gly	Gln	Arg	Ser	Val	Ser	Arg	Arg	Ser
117				325					330					335		
118	Phe	Asn	His	His	Lys	Ala	Arg	Gly	Ala	Gly	Asp	Pro	Lys	Gly	Gln	Arg
119			340					345					350			
120	Trp	His	Thr	Leu	Val	Pro	Leu	Gly	Thr	Glu	Thr	Ile	Thr	Asp	Ser	Tyr
121		355					360					365				
122	Met	Gly	Ala	Pro	Ala	Ser	Glu	Leu	Asp	Thr	Asn	Phe	Phe	Thr	Leu	Tyr
123		370					375				380					
124	Val	Ala	Gln	Gly	Thr	Asn	Lys	Ser	Gln	Gln	Tyr	Lys	Phe	Gly	Thr	Ala
125	385					390					395				400	
126	Thr	Tyr	Ala	Leu	Lys	Glu	Pro	Val	Met	Lys	Ser	Asp	Ser	Trp	Ala	Val
127				405					410					415		
128	Val	Arg	Val	Gln	Ser	Val	Trp	Gln	Leu	Gly	Asn	Arg	Gln	Arg	Pro	Tyr
129			420					425					430			
130	Pro	Trp	Asp	Val	Asn	Trp	Ala	Asn	Ser	Thr	Met	Tyr	Trp	Gly	Gly	Gln
131		435					440					445				
132	Pro	Met	His	Gly	Asn	Asp	Gly	Gln	Pro	Ala	Ala	Gly	Gly	Ser	Glu	Ser
133		450				455					460					
134	Ala	Leu	Ser	Arg	Glu	Gly	Gln	Pro	Gly	Pro	Ser	Gly	Ala	Ala	Gln	Gly
135	465				470					475					480	
136	Gln	Val	Ile	Ser	Asn	Glu	Arg	Ser	Pro	Arg	Arg	Tyr	Ser	Thr	Arg	Thr
137			485						490					495		
138	Ile	Asn	Gly	Val	Gln	Ala	Thr	Asn	Lys	Phe	Thr	Ala	Val	Gly	Asn	Pro
139			500					505					510			
140	Ser	Leu	Gln	Arg	Asp	Pro	Asp	Trp	Tyr	Arg	Trp	Asn	Tyr	Asn	His	Ser
141		515					520					525				
142	Ile	Ala	Val	Trp	Leu	Arg	Glu	Cys	Ser	Arg	Ser	His	Ala	Lys	Ile	Cys
143		530				535					540					
144	Asn	Cys	Gly	Gln	Phe	Arg	Lys	His	Trp	Phe	Gln	Glu	Cys	Ala	Gly	Leu
145	545				550				555						560	
146	Glu	Asp	Arg	Ser	Thr	Gln	Ala	Ser	Leu	Glu	Glu	Ala	Ile	Leu	Arg	Pro
147			565					570					575			
148	Leu	Arg	Val	Gln	Gly	Lys	Arg	Ala	Lys	Arg	Lys	Leu	Asp	Tyr	His	Tyr
149			580					585					590			
150	Ser	Gln	Pro	Thr	Pro	Asn	Arg	Lys	Lys	Val	Tyr	Lys	Thr	Val	Arg	Trp
151		595					600					605				
152	Gln	Asp	Glu	Leu	Ala	Asp	Arg	Glu	Ala	Asp	Phe	Thr	Pro	Ser	Glu	Glu
153		610				615					620					
154	Asp	Gly	Gly	Thr	Thr	Ser	Ser	Asp	Phe	Asp	Glu	Asp	Ile	Asn	Phe	Asp
155	625				630				635						640	

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156 Ile Gly Gly Asp Ser Gly Ile Val Asp Glu Leu Leu Gly Arg Pro Phe
157           645           650           655
158 Thr Thr Pro Ala Pro Val Arg Ile Val Met Asn Ala Leu Gln Glu Asp
159           660           665           670
160 Thr Pro Pro Gly Pro Ser Thr Val Phe Arg Pro Pro Thr Ser Ser Arg
161           675           680           685
162 Pro Leu Glu Thr Pro His Cys Arg Glu Ile Arg Ile Gly Ile Ala Gly
163           690           695           700
164 Ile Thr Ile Thr Leu Ser Leu Cys Gly Cys Ala Asn Ala Arg Ala Pro
165 705           710           715           720
166 Thr Leu Arg Ser Ala Thr Ala Asp Asn Ser Glu Ser Thr Gly Phe Lys
167           725           730           735
168 Asn Val Pro Asp Leu Arg Thr Asp Gln Pro Lys Pro Pro Ser Lys Lys
169           740           745           750
170 Arg Ser Cys Asp Pro Ser Glu Tyr Arg Val Ser Glu Leu Lys Glu Ser
171           755           760           765
172 Leu Ile Thr Thr Thr Pro Ser Arg Pro Arg Thr Ala Arg Arg Cys Ile
173           770           775           780
174 Arg Leu
175 785

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VERIFICATION SUMMARY

PATENT APPLICATION: US/10/591,725

DATE: 09/13/2006

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Input Set : A:\021080002U4.txt

Output Set: N:\CRF4\09132006\J591725.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application Number